

RESEARCH ARTICLE

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Twelve years' detection of respiratory viruses by immunofluorescence in hospitalised children: impact of the introduction of a new respiratory picornavirus assay

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Abstract

Background: Direct immunofluorescence assays (DFA) are a rapid and inexpensive method for the detection of respiratory viruses and may therefore be used for surveillance. Few epidemiological studies have been published based solely on DFA and none included respiratory picornaviruses and human metapneumovirus (hMPV). We wished to evaluate the use of DFA for epidemiological studies with a long-term observation of respiratory viruses that includes both respiratory picornaviruses and hMPV.

Methods: Since 1998 all children hospitalized with respiratory illness at the University Hospital Bern have been screened with DFA for common respiratory viruses including adenovirus, respiratory syncytial virus (RSV), influenza A and B, and parainfluenza virus 1-3. In 2006 assays for respiratory picornaviruses and hMPV were added. Here we describe the epidemiological pattern for these respiratory viruses detected by DFA in 10'629 nasopharyngeal aspirates collected from 8'285 patients during a 12-year period (1998-2010).

Results: Addition of assays for respiratory picornaviruses and hMPV raised the proportion of positive DFA results from 35% to 58% ($p < 0.0001$). Respiratory picornaviruses were the most common viruses detected among patients ≥ 1 year old. The seasonal patterns and age distribution for the studied viruses agreed well with those reported in the literature. In 2010, an hMPV epidemic of unexpected size was observed.

Conclusions: DFA is a valid, rapid, flexible and inexpensive method. The addition of assays for respiratory picornaviruses and hMPV broadens its range of viral detection. DFA is, even in the "PCR era", a particularly adapted method for the long term surveillance of respiratory viruses in a pediatric population.

Background

Respiratory infections are a major cause of morbidity and hospitalizations in children [1], of which a significant proportion are caused by viruses [2,3]. Surveillance of respiratory viruses is important to predict seasonal epidemics, to define patient risk groups and to allocate hospital resources, as well as to describe the burden and characteristics of emerging viruses [4].

In the field of virology, the most commonly used diagnostic methods for virus detection are culture, rapid

culture (such as shell vial assay), direct immunofluorescence staining of clinical specimens (DFA), and PCR. PCR is attractive due to its high sensitivity and broad range of virus detection. PCR-based studies have suggested the important role of respiratory picornaviruses (rhinovirus and enterovirus) as a leading cause of lower respiratory tract infections in children [5], in particular wheezing illnesses such as bronchiolitis [6,7], wheezy bronchitis [8] and asthma exacerbations [9], but also pneumonia [2]. In addition, PCR has allowed for the detection of new respiratory viruses, such as hMPV [10], which has been implicated in upper and lower respiratory tract infections in children [11-13]. It is widely

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believed that nowadays, epidemiological studies on respiratory viruses can only be done with PCR.

However, the high sensitivity of PCR is also a limitation of the technique. A significant proportion of asymptomatic children test positive by PCR to respiratory viruses [14-16], and picornavirus RNA can be detected by PCR up to 5 weeks after an acute infection [17]. Therefore, epidemiological studies based on PCR may overestimate the burden of certain viruses, in particular the common respiratory picornaviruses.

DFA has lower sensitivity than PCR, but this may be an advantage for the detection of clinically relevant infections [18,19]. Moreover, DFA is more rapid and less expensive than PCR and can therefore be used for real-time, routine surveillance of respiratory viruses, which would be difficult by PCR because of the high costs [20,21]. Nevertheless only few epidemiological studies have been published based solely on DFA [22,23], and none of them included testing for respiratory picornaviruses or hMPV, because antibodies for the detection of hMPV have only recently become available and there are no commercial antibodies for the detection of respiratory picornaviruses. Recently, our group reported the validity of immunofluorescence for the detection of picornaviruses directly in respiratory samples using monoclonal antibodies originally designed for the identification of enterovirus in culture [24].

We aimed to evaluate the use of DFA for epidemiological studies of respiratory viruses, now that assays for respiratory picornaviruses and hMPV are available. We performed a retrospective analysis among pediatric patients hospitalized with respiratory tract infections between 1998 and 2010 at the University Hospital Bern. Prospective DFA testing in nasopharyngeal aspirates has been used routinely in this institution for adenovirus (ADV), respiratory syncytial virus (RSV), influenza A and B (IFA and IFB), and parainfluenza 1-3 (PIV 1-3) since 1998, and additionally for hMPV and respiratory picornaviruses since 2006.

Methods

Patient population and sample collection

The study was approved by the Ethics Committee of the University Hospital of Bern in accordance with cantonal ethical regulations (Nr. E 10-01-10). The study included consecutive respiratory tract samples from children under the age of 17 years, who were hospitalized at the Department of Pediatrics, University Hospital Bern, between May 1st 1998 and April 30th 2010. During the entire study period the pediatrics department had the policy of screening children for respiratory viruses if they were hospitalized with a respiratory illness or if they developed respiratory symptoms during their hospital stay.

A total of 12'189 respiratory samples were collected. After exclusion criteria, 10'629 samples remained for the retrospective analysis of DFA results. The exclusion criteria were as follows: samples other than nasopharyngeal aspirates, samples containing less than 20 epithelial cells and samples not tested against the whole viral panel; results of samples from the same patient taken within a time period of 7 days (considered part of the same respiratory episode); results for the month of August 2009, since during this time period practically all respiratory samples were tested by PCR rather than DFA due to the influenza A H1N1 pandemic.

Direct immunofluorescence testing (DFA) for respiratory viruses

All samples were analysed at the Institute for Infectious Diseases, University of Bern. The methods used have previously been described [24].

Between May 1st 1998 and August 31st 2007, the Light Diagnostics Respiratory Viral Screen DFA (Chemicon International, now Millipore) and single fluorescein-conjugated monoclonal antibodies against ADV, RSV, IFA, IFB and PIV 1-3 were used. From September 1st 2007 to April 30th 2010, the D3 Ultra DFA Respiratory Virus Screening & ID Kit (Diagnostic Hybrids) was used for the same viruses. Starting in March 2006, the Light Diagnostics Pan-Enterovirus Reagent "Blend" (Chemicon International/Millipore) was introduced for the detection of respiratory picornaviruses. This assay is formally an indirect immunofluorescence assay, as described elsewhere [24], and does not allow the differentiation between rhinoviruses and enteroviruses. In November 2006, the DFA Metapneumovirus Identification Kit (Diagnostic Hybrids) was added to the screening.

Statistical analysis

An epidemiological year was defined as May 1st to April 30th of the following year. Summer was defined as the months July to September, and winter as January to March. Epidemiological years were designated "odd" if the month of January was in an odd year, and they were labelled "even" if the month of January was in an even year.

All statistical analyses were performed with the GraphPad Prism 5[®] software tool (GraphPad Software, Inc.). Proportions were compared using the chi-square test. Medians were compared with the Kruskal-Wallis test and Dunn's multiple comparison test. A cut-off of $p \leq 0.05$, two tailed, was used for all statistical analyses.

Results

Population

A total of 10'629 samples from 8'285 patients were analysed. The median age of the study population was 11 months (range 0-17 years) and 57.5% were boys.

Rate of viral detection

Before the addition of DFA for picornavirus and hMPV, the rate of viral detection was dominated by the RSV season, with a yearly average rate of 35%, peaks of up to 64% (average 46%) in winter seasons, and troughs as low as 4% (average 16%) during summer time. The addition of DFA for picornaviruses increased the positivity rate, and dampened the seasonal variations. The positivity rate after 2006 was on average 65% in winter and 52% in summer, with a yearly average of 58% (versus 35%; $p < 0.0001$) (Figure 1, Table 1). For comparison, analysis performed on 256 specimens in parallel to DFA screening with the xTag Respiratory Viral Panel (Luminex Molecular Diagnostics) between November 2006 and September 2007 yielded a positivity rate of 78%. The higher detection rate by PCR could mostly be attributed to increased detection of respiratory picornaviruses; 57 from the 78 additional positive results were respiratory picornaviruses (unpublished data).

Pathogens

Respiratory picornaviruses were the most common pathogens detected overall in our study population after the introduction of the assay (27% versus 21% for RSV; $p < 0.0001$) (Table 1). They were present year round, with peaks in the spring and the fall. During the summer time, respiratory picornaviruses also accounted for the majority of viral respiratory infections (Figure 1). Low prevalence of respiratory picornaviruses during winter time coincided with the winter peaks caused by RSV, influenza, or hMPV, and this was the only time during the year when respiratory picornaviruses were not the most commonly detected respiratory virus (Figure 2).

RSV was the second most commonly detected pathogen after picornaviruses (overall prevalence of 21%), but the most prevalent virus during the winter months (Table 1, Figure 2). It manifested a biennial pattern,

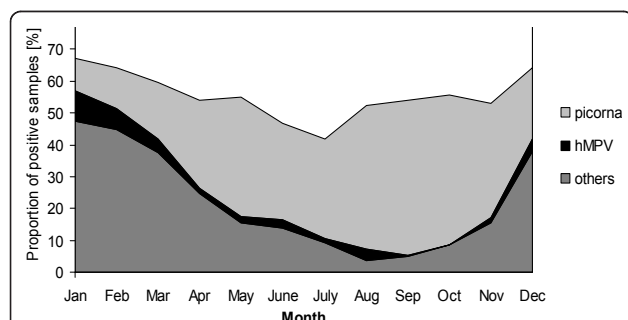


Figure 1 Mean monthly distribution of respiratory viruses detected by DFA in nasopharyngeal aspirates from hospitalized children from November 2006 to April 2010.

picorna = respiratory picornaviruses; hMPV = human metapneumovirus; others = adenovirus, respiratory syncytial virus, influenza A and B, parainfluenza viruses 1-3.

with large winter seasons in odd years alternating with smaller ones in even years.

Influenza A virus was found in 4.1% of all samples and caused yearly winter epidemics, except during the winter season 2005-2006 when it was replaced by influenza B virus (Table 1, Figure 2).

hMPV was detected in 3.4% of all samples collected after November 2006 (Table 1). Yearly hMPV activity varied from being almost absent during the winter season 2006 to 2007, to causing yearly winter outbreaks during the following years (Figure 2, Figure 3). In early 2010, viral activity surpassed previous years, with more cases observed within 5 months (71 cases between December 2009 and April 2010) than during the entire previous period since introduction of the DFA test (56 cases between November 2006 and November 2009).

Age distribution

We compared the proportion of samples positive for a given virus by age (Figure 4). Respiratory picornaviruses were the most common pathogens in children ≥ 1 years (**1-4 years**: 33% versus 15% RSV, $p < 0.0001$; **5-8 years**: 26% versus <7% other viruses, $p < 0.0001$; **9-17 years**: 16% versus 9% influenza A, $p = 0.007$), and RSV was the most common detected in children < 1 year (30% versus 23% picornavirus, $p < 0.0001$). Influenza A showed growing importance with age, and was the second most common virus detected in children > 9 years old (9% influenza A versus <4% other viruses, $p < 0.0001$).

Codetection of respiratory viruses

Out of the total of 10'629 samples analysed, 82 were positive for two viruses (Table 2). No sample was positive for more than two viruses. This corresponds to a codetection rate of 0.8% with DFA.

With the xTag Respiratory Viral Panel (Luminex) 10.2% of samples were positive for two viruses (in 81% of these a respiratory picornavirus was present). In 0.8% of samples we detected three respiratory viruses.

Discussion

In order to determine the value of DFA in conducting epidemiological studies on respiratory viruses now that assays for respiratory picornaviruses and hMPV are available, we retrospectively analysed the results of 12 years of DFA screening of viral pathogens in hospitalized children with respiratory disease.

Respiratory picornaviruses were the most common viral pathogens detected overall in our study, with the exception of patients < 1 year in whom RSV was detected more often, confirming the results of recent studies based on molecular methods [25,26]. PCR detection of respiratory picornaviruses suggests a previously unexpected role [27,28] in severe respiratory disease

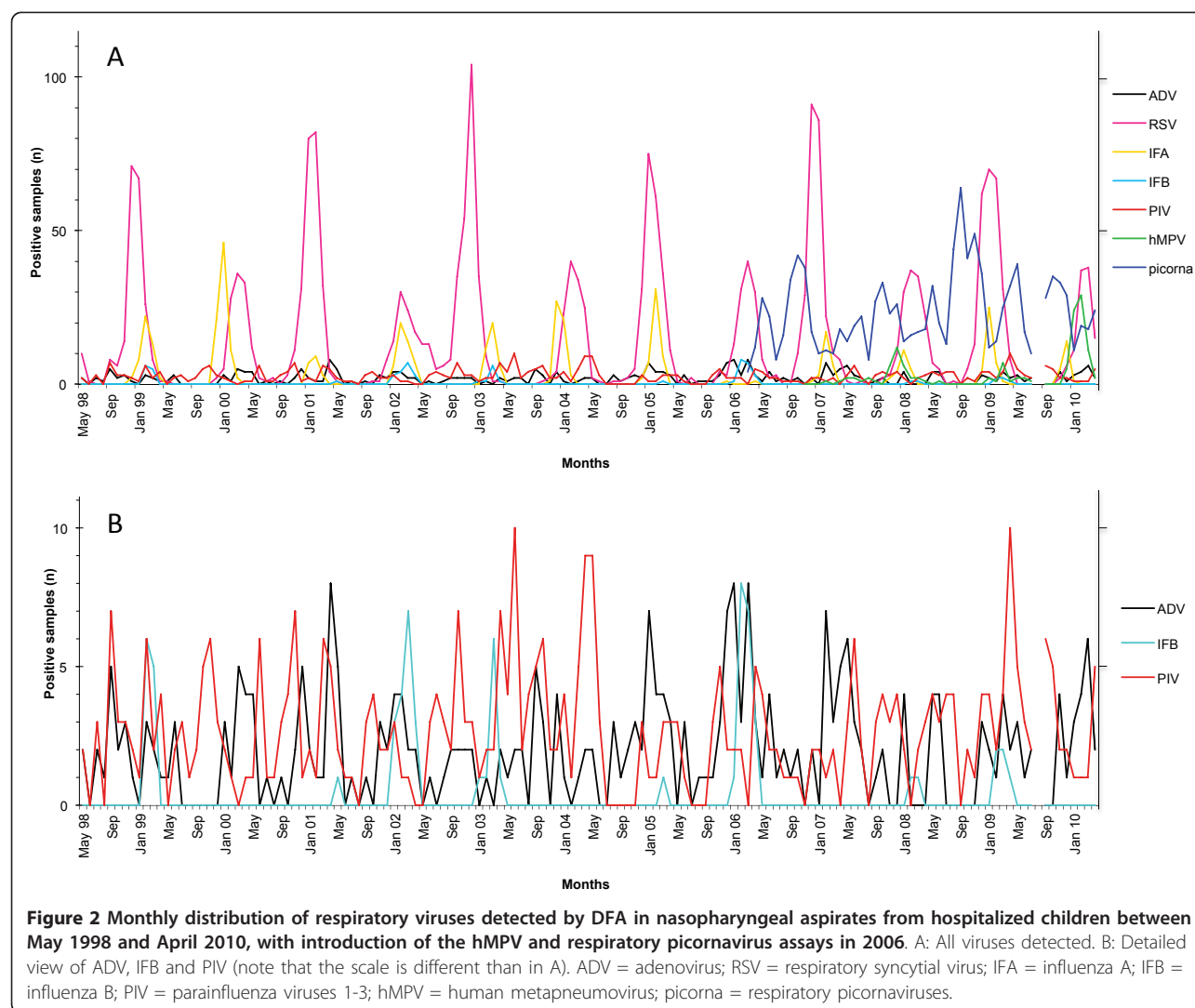
Table 1 Proportion (%) of positive DFA results in nasopharyngeal aspirates by virus and by year, 1998-2010

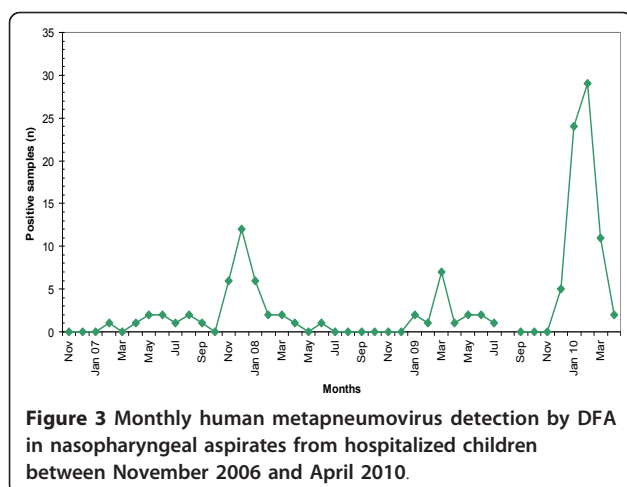
Epidemiological years	98/99	99/00	00/01	01/02	02/03	03/04	04/05	05/06	06/07	07/08	08/09	09/10	1998-2010	1998-2006	2006-2010
ADV	3	3	3	3	1	2	4	4	3	2	2	3	3		
RSV	32	15	32	12	31	15	26	13	26	15	23	12	21		
IFA	7	12	2	6	4	6	6	0.2	3	2	3	2	4		
IFB	2	0	0	2	1	0	0.1	2	0	0.2	0.4	0	0.6		
PIV	5	4	5	3	4	6	3	2	2	3	4	4	4		
hMPV									0.2	4	1	8	3		
Picorna								2	24	26	33	28	27		
Total	49	34	43	26	41	29	38	24	57	53	66	57	44	35	58
Summer	28	16	15	10	24	20	8	4	48	48	63	43		16	52
Winter	58	42	58	34	41	40	64	33	60	60	70	68		46	65

An epidemiological year lasts from May 1st to April 30th of the following year.

ADV = adenovirus; RSV = respiratory syncytial virus; IFA = influenza A; IFB = influenza B; PIV = parainfluenza viruses 1-3;

hMPV = human metapneumovirus; picorna = respiratory picornaviruses.





[25,26,29], but this issue is still debated, given that viral genome can also be detected by PCR many weeks after an acute viral infection [17], or even in entirely asymptomatic children [16,30]. DFA assays need a high viral load to score positive, so a positive result may be more indicative of an acute infection caused by the virus [18,19]. Our high detection rate of respiratory picornaviruses by DFA in hospitalized patients therefore supports their high burden of disease.

The introduction of the hMPV and respiratory picornavirus assays in 2006 increased the positivity rate of our DFA screening from 35% to 58%. For comparison, PCR methods in our laboratory and in the literature usually reach positivity rates of well over 70%, in large part due to a higher detection of respiratory picornaviruses [6,7,26,31]. DFA's lower sensitivity, in particular for respiratory picornaviruses, can however be seen as an advantage considering the difficulty in interpreting

Table 2 Number of nasopharyngeal aspirates with viral codetections by DFA

	picorna	hMPV	PIV	IFB	IFA	RSV
ADV	9	2	4	0	2	10
RSV	29	0	0	1	7	
IFA	1	0	4			
PIV	9	1				
hMPV	3					

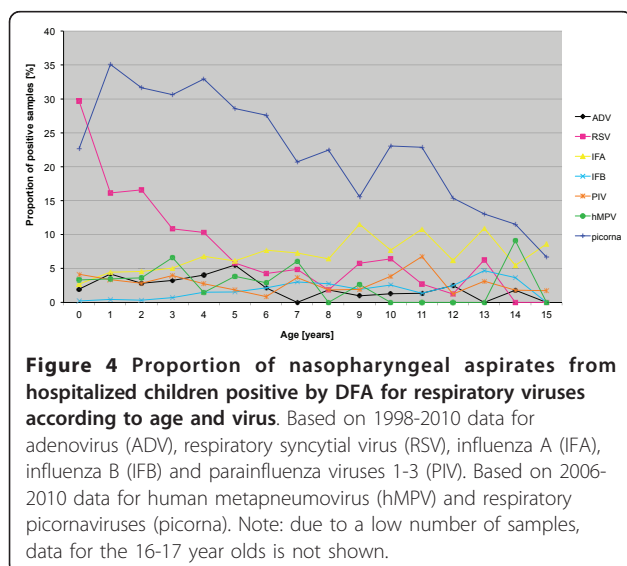
ADV = adenovirus; RSV = respiratory syncytial virus; IFA = influenza A; IFB = influenza B; PIV = parainfluenza viruses 1-3; hMPV = human metapneumovirus; picorna = respiratory picornaviruses.

the clinical significance of PCR-positive results, as described above.

Another common issue in PCR-based studies is the high codetection rate, with on average about 20% of samples being positive for two or more viruses [3,26]. With the xTag Respiratory Viral Panel we detected more than one virus in 11% of samples. In 81% of these a respiratory picornavirus was present. The clinical significance of these "coinfections" remains unclear because of the high sensitivity of PCR [3,32,33], especially for respiratory picornaviruses. It is difficult to determine whether both or only one and which of the codetected pathogens is causing the acute illness [30]. Our lower rate of codetection by DFA suggests that most codetections detected by PCR may indicate consecutive infections.

With the exception of hMPV, our study confirms known patterns of seasonality and age distribution for the studied viruses [22,26,34,35]. It has been postulated that hMPV has a biennial "large-early" and "small-late" season cycle [26,36-38]. We observed an unexpectedly large epidemic in early 2010, which was observed simultaneously in many cities throughout Germany (personal communication, Prof. O. Adams, University of Düsseldorf). Given the recent discovery of hMPV [10], epidemiological studies so far have covered a short time interval and continued monitoring is necessary.

Currently, PCR is considered the most adapted technique to conduct epidemiological studies on respiratory viruses. In contrast to molecular methods, DFA is low in cost and has a rapid turnaround time [21]. Assays can be performed many times a day, and one does not have to wait for a certain number of samples to be collected to start a run. The samples can be screened for many different viruses simultaneously ("multiplex"). Results are usually available within 2-3 hours [39]. These aspects make DFA a method widely and often used in clinical routine, and this concurrently provides the data for ongoing, real-time surveillance of circulating viral pathogens on a large scale. Our systematic monitoring led for example to the early detection of the unexpectedly large hMPV epidemic mentioned above.



Conclusions

In conclusion, DFA's clinical relevance, flexibility and capacity to conduct "multiplex" assays at very low cost make it a valuable diagnostic tool, and now that its range of viral detection has been broadened to include hMPV and especially respiratory picornaviruses, allows for long-term, systematic, real-time monitoring of local epidemiology in pediatric populations.

Abbreviations

DFA: direct immunofluorescence assay; ADV: adenovirus; RSV: respiratory syncytial virus; IFA: influenza A; IFB: influenza B; PIV 1-3: parainfluenza viruses 1-3; hMPV: human metapneumovirus; picorna: respiratory picornaviruses;

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Authors' contributions

CDS participated in statistical analysis, data interpretation, drafting of the manuscript, and critical revision of the manuscript. CA participated in study design and sample acquisition. MGH participated in study design, data interpretation, and critical revision of the manuscript. MTB participated in study design, data interpretation, drafting of the manuscript and critical revision of the manuscript. All authors read and approved the final manuscript.

Competing interests

The authors declare that they have no competing interests.

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